



Fig.22

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1  atgaaataacctgctgccgaccgctgctgctggtctgctgctcctc
_M K Y L L P T A A A G L L L L
46  gctgcccagccggcgatggccatggacatcaagcataatcggccg
A A Q P A M A M D I K H N R P
91  cgcagagactgcgtggcagagggcaaagtgtgtgacctactgtgc
R R D C V A E G K V C D P L C
136 tcctctgggggatgctggggcccaggccctgggtcagtgtgtgtcc
S S G G C W G P G P G Q C L S
181 tgtcgaaattatagccgaggaggtgtctgtgtgacctactgcaac
C R N Y S R G G V C V T H C N
226 tttctgaatggggagcccctcgagcaccaccaccaccactga (SEQ ID NO:17)
F L N G E P L E H H H H H H (SEQ ID NO:14)

```

Fig. 23

ATGGTTTGTGTAGCCAGCTGTCCCCATAACTTTGTGGTGGATCAAACATCCTGTGTCA
GGGCCTGTCCTCCTGACAAGATGGAAGTAGATAAAAATGGGCTCAAGATGTGTGAGC
CTTGTGGGGGACTATGTCCCAAAGCCTGTGAGGGAACAGGCTCTGGGAGCCGCTTCCA
GACTGTGGACTCGAGCAACATTGATGGATTTGTGAACTGCACCAAGATCCTGGGCAAC
CTGGACTTTCTGATCACCGGCCTCAATGGAGACCCCTGGCACAAGATCCCTGCCCTGG
ACCCAGAGAAGCTCAATGTCTTCCGGACAGTACGGGAGATCACAGGTTACCTGAACA
TCCAGTCCTGGCCGCCCCACATGCACAACTTCAGTGTTTTTTTCCAATTTGACAACCATT
GGAGGCAGAAAGCTTGCGGCCGCACTCGAGCACCACCACCACCACTGA

Fig.24(SEQ ID NO :15)

Met VCVASCPHNFVVDQTSCVRACPPDK Met EVDKNGLK Met CEP C
GGLCPKACEGTGSGSRFQTV DSSNIDGFVNCTKILGNLDFLITGL
NGDPWHKIPALDPEKLN VFR TVREITGYLNIQSWPPH Met HNFSV
FSNLTTIGGRSKLAAALEHHHHH

Fig.25(SEQ ID NO :16)